



SEQUENCE LISTING

<110> Baum, Peter
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<120> Novel DNAs and Polypeptides

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<150> 60/107821

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<210> 18

<211> 2173

<212> DNA

<213> Homo sapiens

<400> 18

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 <212> DNA
 <213> Homo sapiens

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 <211> 1669
 <212> DNA
 <213> Homo sapiens

<400> 20

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ccagaaaaga	acaagcttca	tttgtaaaaa	aggaaaacaa	ctcaggcaat	gggggtggct	420
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<211> 1025

<212> DNA

<213> Homo sapiens

<400> 21

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gccgc						1025

<210> 22
 <211> 1039
 <212> DNA
 <213> Homo sapiens

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<210> 23
 <211> 466
 <212> DNA
 <213> Homo sapiens

<400> 23
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<210> 24
 <211> 32
 <212> PRT
 <213> Homo sapiens

<400> 24
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 Arg Ile Asp Leu Cys Val Ser Pro Asn Lys Leu Thr Tyr Ser Pro Lys
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<210> 25
 <211> 98
 <212> PRT
 <213> Homo sapiens

<400> 25
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 20 25 30
 His Ile Thr Asp Thr Gln Phe Lys Lys Gln Asn Ile Thr Ala Pro Ser
 35 40 45
 Arg Ile Phe Leu Gly Ser Leu Pro Ser Leu Leu Thr Pro Asp Tyr Lys
 50 55 60
 Gln Pro Pro Pro Ile Ser Pro Asp Ile Val Leu Tyr Glu Ser Ser Ser
 65 70 75 80
 Ser Gln Met Gly Leu Phe Cys Pro Leu Gly Thr Leu Gly Ser Ile Trp
 85 90 95
 Arg His

<210> 26
 <211> 663
 <212> PRT
 <213> Homo sapiens

<400> 26
 Met Ile Val Gln Met Thr Val Ile Leu Lys Leu Glu Met Pro Gln Asp
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 Ser Leu Ile Leu Glu Lys Ser Gln Asn Trp Ser Ser Gln Lys Met Asp
 20 25 30
 His Ile Leu Ile Cys Cys Val Cys Leu Gly Asp Asn Ser Glu Asp Ala
 35 40 45
 Asp Glu Ile Ile Gln Cys Asp Asn Cys Gly Ile Thr Val His Glu Gly
 50 55 60
 Cys Tyr Gly Val Asp Gly Glu Ser Asp Ser Ile Met Ser Ser Ala Ser
 65 70 75 80
 Glu Asn Ser Thr Glu Pro Trp Phe Cys Asp Ala Cys Lys Cys Gly Val
 85 90 95
 Ser Pro Ser Cys Glu Leu Cys Pro Asn Gln Asp Gly Ile Phe Lys Glu
 100 105 110
 Thr Asp Ala Gly Arg Trp Val His Ile Val Cys Ala Leu Tyr Val Pro
 115 120 125

Gly	Val	Ala	Phe	Gly	Asp	Ile	Asp	Lys	Leu	Arg	Pro	Val	Thr	Leu	Thr	130	135	140	
Glu	Met	Asn	Tyr	Ser	Lys	Tyr	Gly	Ala	Lys	Glu	Cys	Ser	Phe	Cys	Glu	145	150	155	160
Asp	Pro	Arg	Phe	Ala	Arg	Thr	Gly	Val	Cys	Ile	Ser	Cys	Asp	Ala	Gly	165	170	175	
Met	Cys	Arg	Ala	Tyr	Phe	His	Val	Thr	Cys	Ala	Gln	Lys	Glu	Gly	Leu	180	185	190	
Leu	Ser	Glu	Ala	Ala	Ala	Glu	Glu	Asp	Ile	Ala	Asp	Pro	Phe	Phe	Ala	195	200	205	
Tyr	Cys	Lys	Gln	His	Ala	Asp	Arg	Leu	Asp	Arg	Lys	Trp	Lys	Arg	Lys	210	215	220	
Asn	Tyr	Leu	Ala	Leu	Gln	Ser	Tyr	Cys	Lys	Met	Ser	Leu	Gln	Glu	Arg	225	230	235	240
Glu	Lys	Gln	Leu	Ser	Pro	Glu	Ala	Gln	Ala	Arg	Ile	Asn	Ala	Arg	Leu	245	250	255	
Gln	Gln	Tyr	Arg	Ala	Lys	Ala	Glu	Leu	Ala	Arg	Ser	Thr	Arg	Pro	Gln	260	265	270	
Ala	Trp	Val	Pro	Arg	Glu	Lys	Leu	Pro	Arg	Pro	Leu	Thr	Ser	Ser	Ala	275	280	285	
Ser	Ala	Ile	Arg	Lys	Leu	Met	Arg	Lys	Ala	Glu	Leu	Met	Gly	Ile	Ser	290	295	300	
Thr	Asp	Ile	Phe	Pro	Val	Asp	Asn	Ser	Asp	Thr	Ser	Ser	Ser	Val	Asp	305	310	315	320
Gly	Arg	Arg	Lys	His	Lys	Gln	Pro	Ala	Leu	Thr	Ala	Asp	Phe	Val	Asn	325	330	335	
Tyr	Tyr	Phe	Glu	Arg	Asn	Met	Arg	Met	Ile	Gln	Ile	Gln	Glu	Asn	Met	340	345	350	
Ala	Glu	Gln	Lys	Asn	Ile	Lys	Asp	Lys	Leu	Glu	Asn	Glu	Gln	Glu	Lys	355	360	365	
Leu	His	Val	Glu	Tyr	Asn	Lys	Leu	Cys	Glu	Ser	Leu	Glu	Glu	Leu	Gln	370	375	380	
Asn	Leu	Asn	Gly	Lys	Leu	Arg	Ser	Glu	Gly	Gln	Gly	Ile	Trp	Ala	Leu	385	390	395	400
Leu	Gly	Arg	Ile	Thr	Gly	Gln	Lys	Leu	Asn	Ile	Pro	Ala	Ile	Leu	Arg	405	410	415	
Ala	Pro	Lys	Glu	Arg	Lys	Pro	Ser	Lys	Lys	Glu	Gly	Gly	Thr	Gln	Lys	420	425	430	

Thr Ser Thr Leu Pro Ala Val Leu Tyr Ser Cys Gly Ile Cys Lys Lys
 435 440 445

Asn His Asp Gln His Leu Leu Leu Leu Cys Asp Thr Cys Lys Leu His
 450 455 460

Tyr His Leu Gly Cys Leu Asp Pro Pro Leu Thr Arg Met Pro Arg Lys
 465 470 475 480

Thr Lys Asn Ser Tyr Trp Gln Cys Ser Glu Cys Asp Gln Ala Gly Ser
 485 490 495

Ser Asp Met Glu Ala Asp Met Ala Met Glu Thr Leu Pro Asp Gly Thr
 500 505 510

Lys Arg Ser Arg Arg Gln Ile Lys Glu Pro Val Lys Phe Val Pro Gln
 515 520 525

Asp Val Pro Pro Glu Pro Lys Lys Ile Pro Ile Arg Asn Thr Arg Thr
 530 535 540

Arg Gly Arg Lys Arg Ser Phe Val Pro Glu Glu Glu Lys His Glu Glu
 545 550 555 560

Arg Val Pro Arg Glu Arg Arg Gln Arg Gln Ser Val Leu Gln Lys Lys
 565 570 575

Pro Lys Ala Glu Asp Leu Arg Thr Glu Cys Ala Thr Cys Lys Gly Thr
 580 585 590

Gly Asp Asn Glu Asn Leu Val Arg Cys Asp Glu Cys Arg Leu Cys Tyr
 595 600 605

His Phe Gly Cys Leu Asp Pro Pro Leu Lys Lys Ser Pro Lys Gln Thr
 610 615 620

Gly Tyr Gly Trp Ile Cys Gln Glu Cys Asp Ser Ser Ser Ser Lys Glu
 625 630 635 640

Asp Glu Asn Glu Ala Glu Arg Lys Asn Ile Ser Gln Glu Leu Asn Met
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Glu Gln Lys Asn Pro Lys Lys
 660

<210> 27
 <211> 372
 <212> PRT
 <213> Homo sapiens

<400> 27
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Ala Glu Ser Ser Gln Ser Pro Ala Asp Leu Glu Glu Lys Lys Glu Glu

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	50					55					60				
Cys	Leu	Asn	Ser	Leu	Ile	Gln	Val	Phe	Val	Met	Asn	Val	Asp	Phe	Thr
	65					70					75				80
Arg	Ile	Leu	Lys	Arg	Ile	Thr	Val	Pro	Arg	Gly	Ala	Asp	Glu	Gln	Arg
				85					90					95	
Arg	Ser	Val	Pro	Phe	Gln	Met	Leu	Leu	Leu	Leu	Glu	Lys	Met	Gln	Asp
			100					105					110		
Ser	Arg	Gln	Lys	Ala	Val	Arg	Pro	Leu	Glu	Leu	Ala	Tyr	Cys	Leu	Gln
		115					120					125			
Lys	Cys	Asn	Val	Pro	Leu	Phe	Val	Gln	His	Asp	Ala	Ala	Gln	Leu	Tyr
	130					135					140				
Leu	Lys	Leu	Trp	Asn	Leu	Ile	Lys	Asp	Gln	Ile	Thr	Asp	Val	His	Leu
	145					150					155				160
Val	Glu	Arg	Leu	Gln	Ala	Leu	Tyr	Met	Ile	Arg	Val	Lys	Asp	Ser	Leu
				165					170					175	
Ile	Cys	Val	Asp	Cys	Ala	Met	Glu	Ser	Ser	Arg	Asn	Ser	Ser	Met	Leu
			180					185					190		
Thr	Leu	Pro	Leu	Ser	Leu	Phe	Asp	Val	Asp	Ser	Lys	Pro	Leu	Lys	Thr
		195					200					205			
Leu	Glu	Asp	Ala	Leu	His	Cys	Phe	Phe	Gln	Pro	Arg	Glu	Leu	Ser	Ser
	210					215					220				
Lys	Ser	Lys	Cys	Phe	Cys	Glu	Asn	Cys	Gly	Lys	Lys	Thr	Arg	Gly	Lys
	225					230					235				240
Gln	Val	Leu	Lys	Leu	Thr	His	Leu	Pro	Gln	Thr	Leu	Thr	Ile	His	Leu
				245					250					255	
Met	Arg	Phe	Ser	Ile	Arg	Asn	Ser	Gln	Thr	Arg	Lys	Ile	Cys	His	Ser
			260					265					270		
Leu	Tyr	Phe	Pro	Gln	Ser	Leu	Asp	Phe	Ser	Gln	Ile	Leu	Pro	Met	Lys
		275					280					285			
Arg	Glu	Ser	Cys	Asp	Ala	Glu	Glu	Gln	Ser	Gly	Gly	Gln	Tyr	Glu	Leu
	290					295					300				
Phe	Ala	Val	Ile	Ala	His	Val	Gly	Met	Ala	Asp	Ser	Gly	His	Tyr	Cys
	305					310					315				320
Val	Tyr	Ile	Arg	Asn	Ala	Val	Asp	Gly	Lys	Trp	Phe	Cys	Phe	Asn	Asp

Leu Val Arg Val Ala Cys Asp Leu Arg Leu Leu Val Pro Gly His Phe
 100 105 110
 Trp Asn Phe Gly Lys Met Cys Cys Phe Ala Ser Gly Arg Leu Tyr Leu
 115 120 125
 Val Ala Gly Thr Leu Cys Pro Gln His Thr Phe Phe Cys Asp Ser Arg
 130 135 140
 Gln Lys Gly Gln Met Gln Lys Gln Asn Gly Gly Lys Ala Val Asp Glu
 145 150 155 160
 Arg Gln Leu Phe His Gly Thr Ser Ala Ile Phe Val Asp Ala Ile Cys
 165 170 175
 Gln Gln Asn Phe Asp Trp Arg Val Cys Gly Val His Gly Thr Ser Tyr
 180 185 190
 Gly Lys Gly Ser Tyr Phe Ala Arg Asp Ala Ala Tyr Ser His His Tyr
 195 200 205
 Ser Lys Ser Asp Thr Gln Thr His Thr Met Phe Leu Ala Arg Val Leu
 210 215 220
 Val Gly Glu Phe Val Arg Gly Asn Ala Ser Phe Val Arg Pro Pro Ala
 225 230 235 240
 Lys Glu Gly Trp Ser Asn Ala Phe Tyr Asp Ser Cys Val Asn Ser Val
 245 250 255
 Ser Asp Pro Ser Ile Phe Val Ile Phe Glu Lys His Gln Val Tyr Pro
 260 265 270
 Glu Tyr Val Ile Gln Tyr Thr Thr Ser Ser Lys Pro Ser Val Thr Pro
 275 280 285
 Ser Ile Leu Leu Ala Leu Gly Ser Leu Phe Ser Ser Arg Gln
 290 295 300

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 <211> 31
 <212> PRT
 <213> Homo sapiens

<400> 30
 Met Pro Val Tyr Gly Ile Asn Pro His Leu Thr Phe Gln Pro Ala Ser
 1 5 10 15
 Leu Pro Tyr Gly Phe Arg Thr Cys Gln Pro His Asn Ser Leu Lys
 20 25 30

<210> 31
 <211> 95
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<213> Homo sapiens

<400> 31

Met Leu Ile Glu Asp Val Asp Ala Leu Lys Ser Trp Leu Ala Lys Leu
1 5 10 15
Leu Glu Pro Ile Cys Asp Ala Asp Pro Ser Ala Leu Ala Asn Tyr Val
20 25 30
Val Ala Leu Val Lys Lys Asp Lys Pro Glu Lys Glu Leu Lys Ala Phe
35 40 45
Cys Ala Asp Gln Leu Asp Val Phe Leu Gln Lys Glu Thr Ser Gly Phe
50 55 60
Val Asp Lys Leu Phe Glu Ser Leu Tyr Thr Lys Asn Tyr Leu Pro Leu
65 70 75 80
Leu Glu Pro Val Lys Pro Glu Pro Lys Pro Leu Ala Gln Glu Lys
85 90 95

<210> 32

<211> 261

<212> PRT

<213> Homo sapiens

<400> 32

Met Asp Ser Arg His Thr Phe Ala Pro Ala Ala Met Thr Leu Ser Pro
1 5 10 15
Leu Leu Leu Phe Leu Pro Pro Leu Leu Leu Leu Asp Val Pro Thr
20 25 30
Ala Ala Val Gln Ala Ser Pro Leu Gln Ala Leu Asp Phe Phe Gly Asn
35 40 45
Gly Pro Pro Val Asn Tyr Lys Thr Gly Asn Leu Tyr Leu Arg Gly Pro
50 55 60
Leu Lys Lys Ser Asn Ala Pro Leu Val Asn Val Thr Leu Tyr Tyr Glu
65 70 75 80
Ala Leu Cys Gly Gly Cys Arg Ala Phe Leu Ile Arg Glu Leu Phe Pro
85 90 95
Thr Trp Leu Leu Val Met Glu Ile Leu Asn Val Thr Leu Val Pro Tyr
100 105 110
Gly Asn Ala Gln Glu Gln Asn Val Ser Gly Arg Trp Glu Phe Lys Cys
115 120 125
Gln His Gly Glu Glu Glu Cys Lys Phe Asn Lys Val Glu Ala Cys Val
130 135 140
Leu Asp Glu Leu Asp Met Glu Leu Ala Phe Leu Thr Ile Val Cys Met
145 150 155 160

Glu Glu Phe Glu Asp Met Glu Arg Ser Leu Pro Leu Cys Leu Gln Leu
165 170 175

Tyr Ala Pro Gly Leu Ser Pro Asp Thr Ile Met Glu Cys Ala Met Gly
180 185 190

Asp Pro Gly Met Gln Leu Met His Ala Asn Ala Gln Arg Thr Asp Ala
195 200 205

Leu Gln Pro Pro His Glu Tyr Val Pro Trp Val Thr Val Asn Gly Lys
210 215 220

Pro Leu Glu Asp Gln Thr Gln Leu Leu Thr Leu Val Cys Gln Leu Tyr
225 230 235 240

Gln Gly Lys Lys Pro Asp Val Cys Pro Ser Ser Thr Ser Ser Leu Arg
245 250 255

Ser Val Cys Phe Lys
260

<210> 33

<211> 21

<212> PRT

<213> Homo sapiens

<400> 33

Met Pro Gly Tyr Arg His Cys Thr Pro Ala Trp Val Thr Glu Arg Asp
1 5 10 15

Ser Val Ser Glu Lys
20